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APPLICATION NO.	FILING DATE	FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.
10/026,021	12/21/2001	Yasumichi Hitoshi	021044-001210US	6123

20350 7590 03/25/2005

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EXAMINER

YU, MISOOK

ART UNIT PAPER NUMBER

1642

DATE MAILED: 03/25/2005

Please find below and/or attached an Office communication concerning this application or proceeding.

# Office Action Summary

Application No.

10/026,021

Applicant(s)

HITOSHI ET AL.

Examiner

MISOOK YU, Ph.D.

Art Unit

1642

-- The MAILING DATE of this communication appears on the cover sheet with the correspondence address --  
Period for Reply

A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE 3 MONTH(S) FROM THE MAILING DATE OF THIS COMMUNICATION.

- Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication.
- If the period for reply specified above is less than thirty (30) days, a reply within the statutory minimum of thirty (30) days will be considered timely.
- If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication.
- Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133). Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).

## Status

- 1) ☒ Responsive to communication(s) filed on 27 December 2004 and 13 September 2004.
- 2a) ☐ This action is **FINAL**. 2b) ☒ This action is non-final.
- 3) ☐ Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under *Ex parte Quayle*, 1935 C.D. 11, 453 O.G. 213.

## Disposition of Claims

- 4) ☒ Claim(s) 1-50 is/are pending in the application.
- 4a) Of the above claim(s) 1-8, 12-14, 17, 19, 39-50 is/are withdrawn from consideration.
- 5) ☐ Claim(s) \_\_\_\_\_ is/are allowed.
- 6) ☒ Claim(s) 9-11, 15, 16, 18 and 20-38 is/are rejected.
- 7) ☐ Claim(s) \_\_\_\_\_ is/are objected to.
- 8) ☐ Claim(s) \_\_\_\_\_ are subject to restriction and/or election requirement.

## Application Papers

- 9) ☐ The specification is objected to by the Examiner.
- 10) ☐ The drawing(s) filed on \_\_\_\_\_ is/are: a) ☐ accepted or b) ☐ objected to by the Examiner.  
Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).  
Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d).
- 11) ☐ The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152.

## Priority under 35 U.S.C. § 119

- 12) ☐ Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).
- a) ☐ All b) ☐ Some \* c) ☐ None of:
- ☐ Certified copies of the priority documents have been received.
  - ☐ Certified copies of the priority documents have been received in Application No. \_\_\_\_\_.
  - ☐ Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).

\* See the attached detailed Office action for a list of the certified copies not received.

## Attachment(s)

- 1) ☒ Notice of References Cited (PTO-892)
- 2) ☐ Notice of Draftsperson's Patent Drawing Review (PTO-948)
- 3) ☒ Information Disclosure Statement(s) (PTO-1449 or PTO/SB/08)  
Paper No(s)/Mail Date 6/27/02.
- 4) ☐ Interview Summary (PTO-413)  
Paper No(s)/Mail Date. \_\_\_\_\_.
- 5) ☐ Notice of Informal Patent Application (PTO-152)
- 6) ☒ Other: Exhibit A

## **DETAILED ACTION**

### ***Election/Restrictions***

Applicant's election with traverse of group II in the reply filed on 13 September 2004 is acknowledged. The traversal is on the ground(s) that all of the six groups stem from a common concept and theory, and are thus related. As such, examination of all pending claims would not put a serious burden on the examiner. This is not found persuasive because each different inventions use different active ingredients for different effects as explained in the previous Office action, and search of all six different invention put a serious burden on the examiner.

The requirement is still deemed proper and is therefore made FINAL.

Claims 1-8, 39-50 are withdrawn from further consideration pursuant to 37 CFR 1.142(b), as being drawn to a nonelected invention, there being no allowable generic or linking claim. Claims 12-14, 17, 19 are withdrawn from further consideration pursuant to 37 CFR 1.142(b), as being drawn to a nonelected species, there being no allowable generic or linking claim.

Claims 1-50 are pending, and claims 9-11, 15, 16, 18, 20-38 are examined to the extent they are drawn to the elected species of measuring cellular proliferation. The species election requirement of the different cell lines in claim 29, as set forth in page 3 of the Office action mailed on 8/17/2004, is withdrawn and the search is expanded to other cell lines.

### ***Claim Rejections - 35 USC § 112***

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claims 9-11, 15, 16, 18, 20-38 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.

Claims 9-11, 15, 16, 18, 20-38 are rejected under 35 U.S.C. 112, second paragraph, as being incomplete for omitting essential steps, such omission amounting to a gap between the steps. See MPEP § 2172.01. The omitted steps are: a step that accomplishes the purpose stated in the preamble of the claims. In other words, the claims are missing a step of identifying a compound based on the step (ii).

Claim 9 recites ""under stringent conditions" but it is not clear what the metes and bounds are. The specification at page 15, lines 4-7 discloses "The phrase under stringent hybridization conditions refers to conditions under which a probe will hybridize to its target subsequence, typically in a complex mixture of nucleic acids, but to no other sequences. Stringent conditions are sequence-dependent and will be different in different circumstances." This definition does not reasonably set the claimed property boundary by the limitation. Since what will hybridize would be dependent upon to the hybridization conditions being used, the scope of the claimed invention is indefinite. All the dependent claims, except claim 33 drawn to only one species, are also rejected.

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

Claims 9-11, 15, 16, 18, 20-32, and 34-38 are rejected under 35 U.S.C. 112, first paragraph, as failing to comply with the written description requirement. The claim(s) contains subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention.

This rejection is made because the claimed invention is interpreted as drawn to method of identifying a useful compound using a genus of polypeptides encoded by nucleic acids that hybridize under stringent conditions to a nucleic acid encoding SEQ ID NO:2.

The applicable standard for the written description requirement can be found: MPEP 2163; *University of California v. Eli Lilly*, 43 USPQ2d 1398 at 1407; PTO Written Description Guidelines; *Enzo Biochem Inc. v. Gen-Prove Inc.*, 63 USPQ2d 1609; *Vas-Cath Inc. v. Mahurkar*, 19USPQ2d 1111; and *University of Rochester v. G.D. Searle & Co.*, 69 USPQ2d 1886 (CA FC 2004).

To provide adequate written description and evidence of possession of a claimed genus, the specification must provide sufficient distinguishing identifying characteristics of the genus. The factors to be considered include disclosure of complete or partial structure, physical and/or chemical properties, functional characteristics, structure/function correlation, methods of making the claimed product, or any combination thereof. In this case, the only factor present in the claims is a partial structure in the form of hybridization under the undefined conditions. The claims do not define any function associated with the claimed genus. Accordingly, in the absence of

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sufficient recitation of distinguishing identifying characteristics, the specification does not provide adequate written description of the claimed genus.

Claims 9, and 35 are rejected under 35 U.S.C. 112, first paragraph, as failing to comply with the enablement requirement. The claim(s) contains subject matter which was not described in the specification in such a way as to enable one skilled in the art to which it pertains, or with which it is most nearly connected, to make and/or use the invention.

This rejection is made because the Office interprets the claimed invention as drawn to method of identifying an antisense that modulates cellular proliferation comprising the steps of contacting an antisense to a SAK polypeptide and determining the functional effect of the antisense upon the polypeptide.

The factors considered when determining if the disclosure satisfies the enablement requirement and whether any necessary experimentation is Aundue≡ include, but are not limited to: 1) nature of the invention, 2) state of the prior art, 3) relative skill of those in the art, 4) level of predictability in the art, 5) existence of working examples, 6) breadth of claims, 7) amount of direction or guidance by the inventor, and 8) quantity of experimentation needed to make or use the invention. *In re Wands*, 858 F.2d 731, 737, 8 USPQ2d 1400, 1404 (Fed. Cir. 1988).

The specification does not teach any method of screening an antisense by contacting an antisense to a SAK protein. Antisense is defined as ""having a complementary sequence to a segment of genetic material (as mRNA) and serving to

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inhibit gene function" by Merriam-Webster Online dictionary downed on 3-18-2005 from the url...<http://www.m-w.com/cgi-bin/dictionary?book=Dictionary&va=antisense>. This definition indicates that an antisense works at the nucleic acid expression level. In other words, contacting an antisense to "a SAK polypeptide" would not have any effect. The Office cites US 5,650,501 A (22 July 1997) to show the state of art involving an antisense. The '501 patent teaches at columns 26-27 (Example 4) that a method of transfecting an antisense in order to inhibit an SAK encoding gene expression. The '501 patent teaches an antisense inhibits nucleic acid (gene) expression.

Considering the unpredictable state of art, no guidance, no examples in the specification how to use the instantly claimed invention, broad breath of the claims, it is concluded that undue experimentation is required to practice the invention.

### ***Claim Rejections - 35 USC § 102***

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless –

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

Claims 9-11, 15, 16, 18, 24-32, 34, 36, and 37 are rejected under 35

U.S.C. 102(b) as being anticipated by US 5,650,501 A (IDS AA filed on 06/27/002, 22 July 1997, the '501 patent from now on).

Claims 9-11, 15, 16, 18, 20, 24-32, 34, 36, and 37 are broadly interpreted as drawn to method of identifying a useful compound by determining the phenotypic effect of said compound in cellular proliferation when said compound is contacted with a SAK

polypeptide encoded by a nucleic acid hybridizes under stringent conditions to the nucleic acid encoding the instant SEQ ID NO:2 protein.

The '501 patent teaches SEQ ID NO:3 (a human cDNA) encoding SEQ ID NO:4 (a human SAK protein) that meets the instantly claimed limitation of "a SAK polypeptide encoded by a nucleic acid hybridizes under stringent conditions to the nucleic acid encoding the instant SEQ ID NO:2 protein". Note that the attached sequence alignment (Exhibit A) showing that instant SEQ ID NO:2 and the N-terminal half of SEQ ID NO:4 of the '501 patent have about 90 % homology in protein level, and instant SEQ ID NO:1 and SEQ ID NO:3 of the '501 patent have about 80.9 % local similarity. The Office does not have the facilities and resources to provide the factual evidence needed in order to establish that SEQ ID NO:3 of the '501 patent does not hybridize under the indefinite hybridization conditions to the instant SEQ ID NO:1. In the absence of evidence to the contrary, the burden is on the applicant to prove that the claimed nucleic acid is different from those taught by the prior art and to establish patentable differences. See *In re Best* 562 F.2d 1252, 195 USPQ 430 (CCPA 1977) and *Ex parte Gray* 10 USPQ 2d 1922 (PTO Bd. Pat. App. & Int. 1989).

As for the contacting a compound the polypeptide, the '501 patent discloses at the line bridging columns 1 and 2 that an antisense to block the expression of SAK inhibits cellular proliferation, i.e. "cell growth was suppressed", and at column 5 lines 5-40 discloses "the method comprises providing a known concentration of a serine/threonine kinase protein of the invention, or an isoform or part of the protein, incubating the kinase protein, isoform or part of the protein with a substance which is a



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substrate of the kinase protein, or isoform or part of the protein, and a suspected agonist or antagonist substance, under conditions which permit the phosphorylation of the substrate, and assaying for phosphorylation of the substrate. In a second embodiment, the method comprises providing a known concentration of a serine/threonine kinase protein of the invention, or an isoform or part of the protein, incubating the kinase protein with a substance which is capable of binding to and activating the kinase protein, or isoform or part of the protein, and a suspected agonist or antagonist substance under conditions which permit the formation of substance-protein complexes, and assaying for activation of the kinase protein. The methods of the invention permit the identification of potential stimulators or inhibitors of cell proliferation which will be useful in the treatment of proliferative disorders.” In other words, the invention is to discover the antagonist or agonist of cellular proliferation modulated by the activity of SAK polypeptide.

Further, the '501 patent at column 5 line 23-25 teaches “Substance which affect cell proliferation may be identified”, and “The invention provides a method for screening for substances having pharmaceutical utility in treatment and diagnosis of proliferative disorders”. The '501 patent at column 14 teaches an antibody, method of using the antibody in determining cellular proliferation modulation at column 16, detailed screening assays for measuring cellular proliferation using the SAK polypeptides and other putative medically useful compounds of peptide and antibody from columns 17-20.

As for instant claim 25, and the various cancer cells in claim 26-27, and cancer cells with p53 status of being wild type, the null, or mutant, the '501 patent teaches at

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column 19 lines 52-67 "Substances which are capable of binding to the kinase protein of the invention or isoforms or parts thereof, particularly regulators, agonists and antagonists of the binding of regulators and substrates of Sak protein identified by the methods of the invention, antisense nucleic acid molecules of the invention, and antibodies of the invention may be used for stimulating or inhibiting cell proliferation. The regulators, agonists and antagonists, substrates etc. may accordingly be used to stimulate or inhibit cell proliferation associated with disorders including various forms of cancer such as leukemias, lymphomas (Hodgkins and non-Hodgkins), sarcomas, melanomas, adenomas, carcinomas of solid tissue, hypoxic tumors, squamous cell carcinomas of the mouth, throat, larynx, and lung, genitourinary cancers such as cervical and bladder cancer, hematopoietic cancers, head and neck cancers, and nervous system cancers". Since the instant specification is not about which cancer has null, or mutation, or wild-type in p53, it is the Office's position that various cancer cells of the '501 patent have the different status in p53 gene. The Office does not have the facilities and resources to provide the factual evidence needed in order to establish that the various cancers of the '501 patent do not have the three different p53 status. This determination requires sequencing of all the cancers listed in the '501 patent. In the absence of evidence to the contrary, the burden is on the applicant to prove that the claimed cancer cells are different from those taught by the prior art and to establish patentable differences. See *In re Best* 562 F.2d 1252, 195 USPQ 430 (CCPA 1977) and *Ex parte Gray* 10 USPQ 2d 1922 (PTO Bd. Pat. App. & Int. 1989).

As for claims 30, and 29, drawn to transformed cancer cell lines, the '501 patent at column 12, line 7 teaches "HeLa" cell.

***Claim Rejections - 35 USC § 103***

The following is a quotation of 35 U.S.C. 103(a) which forms the basis for all obviousness rejections set forth in this Office action:

(a) A patent may not be obtained though the invention is not identically disclosed or described as set forth in section 102 of this title, if the differences between the subject matter sought to be patented and the prior art are such that the subject matter as a whole would have been obvious at the time the invention was made to a person having ordinary skill in the art to which said subject matter pertains. Patentability shall not be negated by the manner in which the invention was made.

Claims 9, 15, and 20-23 are rejected under 35 U.S.C. 103(a) as being unpatentable over US 5,650,501 A (22 July 1997) in view of US 5,959,081 A (28 September 1999, the '081 patent from now on).

Claims 9, 15, and 20-23 are interpreted as drawn to method of identifying a useful compound by determining whether or not said compound modulates cellular proliferation, when said compound is contacted with a SAK polypeptide encoded by a nucleic acid hybridizes under stringent conditions to the nucleic acid encoding the instant SEQ ID NO:2 protein, wherein said cellular proliferation is determined by measuring DNA synthesis or measuring green fluorescent protein.

See 102(b) rejection above for what the '501 patent teaches.

The '501 patent does not teach measuring DNA synthesis as an amount of <sup>3</sup>H thymidine incorporation or measuring green fluorescent protein.

However, the '801 patent teaches at columns 24 and 26 that DNA synthesis as an amount of <sup>3</sup>H thymidine incorporation or measuring green fluorescent protein

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detection are well known techniques in the art before the effective filing date of the instant application.

Therefore, it would have been obvious to one of ordinary skill in the art to use DNA synthesis as an amount of  $^3\text{H}$  thymidine incorporation or measuring green fluorescent protein detection with a reasonable expectation of success, given that the '501 patent teaches that a SAK protein is involved in cellular proliferation. One of ordinary skill would be motivated to identify a compound that dilutes the green emission as an candidate that might be inhibiting cellular protein, or the compound that inhibits  $^3\text{H}$  thymidine incorporation in DNA of the cell as a possible candidate for inhibiting cancer cell growth, given that the '501 patent teaches that a SAK protein is involved in cellular proliferation

Claims 9, 37, and 38 are rejected under 35 U.S.C. 103(a) as being unpatentable over US 5,650,501 A (22 July 1997) in view of US 5,589,356 A (31 December 1996, the '356 patent from now on).

Claims 9, 37, and 38 are interpreted as drawn to method of identifying a useful circular peptide by determining whether or not said circular peptide affecting cellular proliferation when said compound is contacted with a SAK polypeptide encoded by a nucleic acid hybridizes under stringent conditions to the nucleic acid encoding the instant SEQ ID NO:2 protein.

See 102(b) rejection above for what the '501 patent teaches.

The '501 patent does not teach a circular peptide.

However, the '356 patent teaches (at the front page) a circular peptide and also teach that a usefulness of a circular peptide as a therapeutic has been recognized in the art before the effective filing date of the instant application (note column 3, lines 3-4).

Therefore, it would have been obvious to one of ordinary skill in the art to add a circular peptide to see whether the circular peptide modulates cellular proliferation, given that the '501 patent teaches that a SAK protein is involved in cellular proliferation and the '356 patent teaches many circular peptides. One of ordinary skill would have been motivated to screen a circular peptide with the art-known detection methods as described by the '501 patent, given that the '356 patent teaches that a circular peptide might be a candidate therapeutic.

### ***Conclusion***

Any inquiry concerning this communication or earlier communications from the examiner should be directed to MISOOK YU, Ph.D. whose telephone number is 571-272-0839. The examiner can normally be reached on 8 A.M. to 5:30 P.M., every other Friday off.

If attempts to reach the examiner by telephone are unsuccessful, the examiner's supervisor, Jeffrey Siew can be reached on 571-272-0787. The fax phone number for the organization where this application or proceeding is assigned is 703-872-9306. Any inquiry of a general nature, matching or filed papers or relating to the status of this application or proceeding should be directed to the Judy Ladrangan for Art Unit 1642 whose telephone number is 571-272-0536.

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Information regarding the status of an application may be obtained from the Patent Application Information Retrieval (PAIR) system. Status information for published applications may be obtained from either Private PAIR or Public PAIR. Status information for unpublished applications is available through Private PAIR only. For more information about the PAIR system, see <http://pair-direct.uspto.gov>. Should you have questions on access to the Private PAIR system, contact the Electronic Business Center (EBC) at 866-217-9197 (toll-free).

MISOOK YU, Ph.D.  
Examiner  
Art Unit 1642

A handwritten signature in black ink, appearing to read "Misook Yu", with a stylized flourish at the end.

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OM protein - protein search, using sw model

Run on: September 23, 2004, 20:59:21 / Search time 22 Seconds  
(without alignments)  
2276.236 Million cell updates/sec

Title: US-10-026-021-2

Perfect score: 5078  
Sequence: 1 MATCIQKIEIDFKVGNLIGK.....KIQCTSLILMFSPNPNH 970

Scoring table: BLOSUM62  
Gapop 10.0, Gapext 0.5

Searched: 389414 seqs, 51625971 residues

Total number of hits satisfying chosen parameters: 389414

Minimum DB seq length: 0  
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%  
Maximum Match 100%

Listing first 45 summaries

Database : Issued Patents AA:\*  
1: /cgn2\_6/prodata/2/1aa/5A COMB pep:\*  
2: /cgn2\_6/prodata/2/1aa/5B COMB pep:\*  
3: /cgn2\_6/prodata/2/1aa/6A COMB pep:\*  
4: /cgn2\_6/prodata/2/1aa/6B COMB pep:\*  
5: /cgn2\_6/prodata/2/1aa/6C COMB pep:\*  
6: /cgn2\_6/prodata/2/1aa/6D COMB pep:\*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	3927.5	77.3	925	1 US-08-252-995D-4	Sequence 4, Appl1
2	3927.5	77.3	925	2 US-08-834-108-4	Sequence 4, Appl1
3	1926	37.9	464	1 US-08-252-995D-6	Sequence 6, Appl1
4	1926	37.9	464	2 US-08-834-108-6	Sequence 6, Appl1
5	1883.5	37.1	416	1 US-08-252-995D-2	Sequence 2, Appl1
6	1883.5	37.1	416	2 US-08-834-108-2	Sequence 2, Appl1
7	1370	27.0	273	1 US-08-252-995D-10	Sequence 10, Appl1
8	1370	27.0	273	2 US-08-834-108-10	Sequence 10, Appl1
9	590.5	11.6	607	3 US-08-272-796-15	Sequence 15, Appl1
10	590.5	11.6	607	2 US-08-834-108-15	Sequence 15, Appl1
11	578.5	11.4	271	1 US-08-252-995D-11	Sequence 11, Appl1
12	578.5	11.4	271	2 US-08-834-108-11	Sequence 11, Appl1
13	561.5	11.1	272	1 US-08-252-995D-12	Sequence 12, Appl1
14	561.5	11.1	272	2 US-08-834-108-12	Sequence 12, Appl1
15	560.5	11.0	685	3 US-08-788-989-1	Sequence 1, Appl1
16	560.5	11.0	685	2 US-08-136-282-2	Sequence 2, Appl1
17	560.5	11.0	685	3 US-09-372-796-1	Sequence 1, Appl1
18	560.5	11.0	685	2 US-09-505-744-2	Sequence 2, Appl1
19	545	10.7	603	4 US-09-311-311C-26	Sequence 26, Appl1
20	533	10.5	603	3 US-09-198-112-2	Sequence 2, Appl1
21	525.5	10.3	272	1 US-08-252-995D-14	Sequence 14, Appl1
22	525.5	10.3	272	2 US-08-834-108-14	Sequence 14, Appl1
23	508.5	10.0	403	2 US-08-755-728-4	Sequence 4, Appl1
24	508.5	10.0	403	1 US-08-974-655-4	Sequence 4, Appl1
25	508.5	10.0	403	3 US-09-283-011-4	Sequence 4, Appl1
26	498	9.8	722	4 US-08-817-832B-32	Sequence 32, Appl1
27	495.5	9.8	722	4 US-09-984-890-4	Sequence 4, Appl1

28	490	9.6	275	1 US-08-252-995D-13	Sequence 13, Appl1
29	490	9.6	275	2 US-08-834-108-13	Sequence 13, Appl1
30	487	9.6	724	4 US-09-984-890-2	Sequence 2, Appl1
31	485.5	9.6	344	2 US-08-755-728-3	Sequence 3, Appl1
32	485.5	9.6	344	2 US-08-974-655-3	Sequence 3, Appl1
33	485.5	9.6	344	3 US-09-283-011-3	Sequence 3, Appl1
34	480	9.5	347	2 US-09-16-000-1	Sequence 1, Appl1
35	479.5	9.4	745	4 US-09-523-849-36	Sequence 36, Appl1
36	464	9.1	729	2 US-08-677-298-2	Sequence 2, Appl1
37	464	9.1	729	4 US-09-523-849-33	Sequence 33, Appl1
38	462.5	9.1	633	3 US-08-557-006C-43	Sequence 43, Appl1
39	454.5	9.0	556	2 US-09-016-000-4	Sequence 4, Appl1
40	454.5	9.0	556	4 US-09-156-792D-2	Sequence 2, Appl1
41	450	8.9	1037	4 US-09-428-711A-21	Sequence 21, Appl1
42	449	8.8	793	4 US-09-523-849-32	Sequence 32, Appl1
43	446.5	8.8	556	4 US-09-800-960-4	Sequence 4, Appl1
44	446.5	8.8	556	4 US-10-096-960-4	Sequence 4, Appl1
45	445.5	8.8	1203	4 US-09-799-875-5	Sequence 5, Appl1

ALIGNMENTS

RESULT 1  
US-08-252-995D-4  
Sequence 4, Application US/08252995D  
Patent No. 5650501  
GENERAL INFORMATION:  
APPLICANT: Dennis, James W  
APPLICANT: Heffernan, Mike  
TITLE OF INVENTION: NOVEL SERINE/THREONINE KINASE  
NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: BERSKIN & PARR  
STREET: 40 King Street West  
CITY: Toronto  
STATE: Ontario  
COUNTRY: Canada  
ZIP: M5H 3T2  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/252,995D  
FILING DATE: 02-JUN-1994  
CLASSIFICATION: 536  
ATTORNEY/AGENT INFORMATION:  
NAME: Kirdydyk, Linda W  
REGISTRATION NUMBER: 34,971  
REFERENCE/DOCKET NUMBER: 3153-96  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (416) 364-7311  
TELEFAX: (416) 361-1398  
INFORMATION FOR SEQ ID NO: 4:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 925 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-252-995D-4  
Query Match 77.3% Score 3927.5; DB 1; Length 925;  
Best Local Similarity 78.6% Pred. No. 8.4e-296;  
Matches 763; Conservative 76; Mismatches 83; Indels 49; Gaps 9;  
QY 1 MATCIQKIEIDFKVGNLIGKSPAGVTRAESIHTEVAIAIKIDKANKYKAGVORVONE 60  
DB 1 MAACIGRIEIDFKVGNLIGKSPAGVTRAESIHTEVAIAIKIDKANKYKAGVORVONE 60  
QY 61 VKIHQQLGHPISLIELVYVFEDSNVYVLLVLEHCHNGENNRILKNRYKPSBNEARHPHQI 120

61 VKHQQLHPSVLEIYNYFEDNNVYVYLEMCHNGEMNYYLKNRMKPFSEBBAHPHQI 120  
121 ITGMLYHSHGILHRDLTSLNLLTRNNMIKIADFGIATOLKMPHEKTYTLCTPNYISP 180  
121 ITGMLYHSHGILHRDLTSLNLLTRNNMIKIADFGIATOLKMPHEKTYTLCTPNYISP 180  
181 EIATRSAGLESDDWSLGCMPYTLIGRPEDTDVKNYLVADYEMPSLEAKD 240  
181 EIATRSAGLESDDWSLGCMPYTLIGRPEDTDVKNYLVADYEMPSLEAKD 240  
241 LIHOLLERNADRLSLSSVLDHPMSRNSSTKSDLGTVEDSIDGATISTATASSST 300  
241 LIHOLLERNADRLSLSSVLDHPMSRNSSTKSDLGTVEDSIDGATISTATASSST 300  
301 SLSGSLD-RRLVGGPLPKKIIVFOKNKNSDP--SSGDSNFTCTOWGNPBOEANSRG 358  
301 SLSGSLD-RRLVGGPLPKKIIVFOKNKNSDP--SSGDSNFTCTOWGNPBOEANSRG 358  
359 RVIQDAERPHSRYLRAVSSDRSGTNSQOAKTYTMRCHSAEMLSVKSGGGENE 418  
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419 RYSPDNNANNINPFKEKTSSSSGSFERPDNNQALSNHLCPEKTPFPADPTPOTETVQ 478  
419 RYSPDNNANNINPFKEKTSSSSGSFERPDNNQALSNHLCPEKTPFPADPTPOTETVQ 478  
479 WFGNLOINAHLEKTEYDISPNRDFOGHPDLQDXTSKAATVTKYKNSDASDNASV 538  
479 WFGNLOINAHLEKTEYDISPNRDFOGHPDLQDXTSKAATVTKYKNSDASDNASV 538  
443 WFGNLOINAHLEKTEYDISPNRDFOGHPDLQDXTSKAATVTKYKNSDASDNASV 501  
539 QONTMKTMTALHSKEPIIQOECVFGSDPLSEOSKTRGMEPMGYONRLTSLTSLVAR 598  
539 QONTMKTMTALHSKEPIIQOECVFGSDPLSEOSKTRGMEPMGYONRLTSLTSLVAR 598  
502 QLSANMKNMAHHPKPEVMPQEP--GLHPSEOSKTRGMEPMGYONRLTSLTSLVAR 598  
502 QLSANMKNMAHHPKPEVMPQEP--GLHPSEOSKTRGMEPMGYONRLTSLTSLVAR 598  
599 LKPIROKTKGAVVSLIDSEECVCLVKEVTAQETVETQISSDGTITITYPNGRG 658  
599 LKPIROKTKGAVVSLIDSEECVCLVKEVTAQETVETQISSDGTITITYPNGRG 658  
560 LKPIROKTKGAVVSLIDSEECVCLVKEVTAQETVETQISSDGTITITYPNGRG 619  
560 LKPIROKTKGAVVSLIDSEECVCLVKEVTAQETVETQISSDGTITITYPNGRG 619  
659 LADRPSPDNTISRYSPDLPEKTYMKYGYASRFVULVSKSPKTYTFRYAKTLMENS 718  
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719 PGADPEWPFYDGVKHKTEDFIQVIEKTKSYTLKSESEVNSLKEEIKMYMDHANEHRI 778  
719 PGADPEWPFYDGVKHKTEDFIQVIEKTKSYTLKSESEVNSLKEEIKMYMDHANEHRI 778  
680 PGADPEWPFYDGVKHKTEDFIQVIEKTKSYTLKSESEVNSLKEEIKMYMDHANEHRI 739  
680 PGADPEWPFYDGVKHKTEDFIQVIEKTKSYTLKSESEVNSLKEEIKMYMDHANEHRI 739  
779 CLAESIISSEERKTRSAFPPIIIGRKPGSTSPKALSPSPVDSNYPTDRASFPNMY 838  
779 CLAESIISSEERKTRSAFPPIIIGRKPGSTSPKALSPSPVDSNYPTDRASFPNMY 838  
740 CLAESIISSEERKTRSAFPPIIIGRKPGSTSPKALSPSPVDSNYPTDRASFPNMY 798  
740 CLAESIISSEERKTRSAFPPIIIGRKPGSTSPKALSPSPVDSNYPTDRASFPNMY 798  
839 MHSAPSPDADILNPSMTNEGGLTTPASGDTISSNLSKOCPLPSAOLLSVYKNGW 898  
839 MHSAPSPDADILNPSMTNEGGLTTPASGDTISSNLSKOCPLPSAOLLSVYKNGW 898  
799 VNSAAPSPDADILNPSMTNEGGLTTPASGDTISSNLSKOCPLPSAOLLSVYKNGW 853  
799 VNSAAPSPDADILNPSMTNEGGLTTPASGDTISSNLSKOCPLPSAOLLSVYKNGW 853  
899 ATQUTSAGVWQVNDGSQLVQAGVSSISYTSPPNGQTRYGENEKLPYIKOKLQCLSSI 958  
899 ATQUTSAGVWQVNDGSQLVQAGVSSISYTSPPNGQTRYGENEKLPYIKOKLQCLSSI 958  
854 ATQUTSAGVWQVNDGSQLVQAGVSSISYTSPPNGQTRYGENEKLPYIKOKLQCLSSI 913  
854 ATQUTSAGVWQVNDGSQLVQAGVSSISYTSPPNGQTRYGENEKLPYIKOKLQCLSSI 913  
959 LMFNSNTPNF 969  
959 LMFNSNTPNF 924  
914 LMFNSNTPNF 924

RESULT 2  
US-08-834-108-4  
Sequence 4, Application US/08834108  
Patent No. 5976893

GENERAL INFORMATION:  
APPLICANT: Dennis, James W  
APPLICANT: Hefeffman, Mike  
APPLICANT: Fode, Carol  
TITLE OF INVENTION: NOVEL SERINE/THREONINE KINASE

NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESSEE: BERSKIN & PARR  
STREET: 40 King Street West  
CITY: Toronto  
STATE: Ontario  
COUNTRY: Canada  
ZIP: M5H 3Y2  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
COMPUTER: IBM PC compatible  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent In Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/834,108  
FILING DATE:  
CLASSIFICATION: 536  
ATTORNEY/AGENT INFORMATION:  
NAME: Kurdyak, Linda M  
REGISTRATION NUMBER: 34,971  
REFERENCE/DOCKET NUMBER: 3153-210  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (416) 364-7311  
FAX: (416) 361-1398  
INFORMATION FOR SEQ ID NO: 4:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 925 amino acids  
TYPE: amino acid  
TOPOLOGY: linear  
MOLECULE TYPE: protein  
US-08-834-108-4  
Query Match  
Best Local Similarity 77.3%; Score 3927.5; DB 2; Length 925;  
Best Local Similarity 78.6%; Pred No. 8.4e-296;  
Matches 763; Conservative 76; Mismatches 83; Indels 49; Gaps 9;  
1 MATIGKIEDPKYVNLGKSPAGVYRAESIHGELFVAIKMIDKAYKAGVORVNE 60  
1 MAACIGRIEDPKYVNLGKSPAGVYRAESIHGELFVAIKMIDKAYKAGVORVNE 60  
61 VKHQQLHPSVLEIYNYFEDNNVYVYLEMCHNGEMNYYLKNRMKPFSEBBAHPHQI 120  
61 VKHQQLHPSVLEIYNYFEDNNVYVYLEMCHNGEMNYYLKNRMKPFSEBBAHPHQI 120  
121 ITGMLYHSHGILHRDLTSLNLLTRNNMIKIADFGIATOLKMPHEKTYTLCTPNYISP 180  
121 ITGMLYHSHGILHRDLTSLNLLTRNNMIKIADFGIATOLKMPHEKTYTLCTPNYISP 180  
121 ITGMLYHSHGILHRDLTSLNLLTRNNMIKIADFGIATOLKMPHEKTYTLCTPNYISP 180  
181 EIATRSAGLESDDWSLGCMPYTLIGRPEDTDVKNYLVADYEMPSLEAKD 240  
181 EIATRSAGLESDDWSLGCMPYTLIGRPEDTDVKNYLVADYEMPSLEAKD 240  
181 EIATRSAGLESDDWSLGCMPYTLIGRPEDTDVKNYLVADYEMPSLEAKD 240  
241 LIHOLLERNADRLSLSSVLDHPMSRNSSTKSDLGTVEDSIDGATISTATASSST 300  
241 LIHOLLERNADRLSLSSVLDHPMSRNSSTKSDLGTVEDSIDGATISTATASSST 300  
241 LIHOLLERNADRLSLSSVLDHPMSRNSSTKSDLGTVEDSIDGATISTATASSST 300  
301 SLSGSLD-RRLVGGPLPKKIIVFOKNKNSDP--SSGDSNFTCTOWGNPBOEANSRG 358  
301 SLSGSLD-RRLVGGPLPKKIIVFOKNKNSDP--SSGDSNFTCTOWGNPBOEANSRG 358  
359 RVIQDAERPHSRYLRAVSSDRSGTNSQOAKTYTMRCHSAEMLSVKSGGGENE 418  
359 RVIQDAERPHSRYLRAVSSDRSGTNSQOAKTYTMRCHSAEMLSVKSGGGENE 418  
419 RYSPDNNANNINPFKEKTSSSSGSFERPDNNQALSNHLCPEKTPFPADPTPOTETVQ 478  
419 RYSPDNNANNINPFKEKTSSSSGSFERPDNNQALSNHLCPEKTPFPADPTPOTETVQ 478  
479 WFGNLOINAHLEKTEYDISPNRDFOGHPDLQDXTSKAATVTKYKNSDASDNASV 538  
479 WFGNLOINAHLEKTEYDISPNRDFOGHPDLQDXTSKAATVTKYKNSDASDNASV 538  
443 WFGNLOINAHLEKTEYDISPNRDFOGHPDLQDXTSKAATVTKYKNSDASDNASV 501  
539 QONTMKTMTALHSKEPIIQOECVFGSDPLSEOSKTRGMEPMGYONRLTSLTSLVAR 598  
539 QONTMKTMTALHSKEPIIQOECVFGSDPLSEOSKTRGMEPMGYONRLTSLTSLVAR 598



3 075

2554 CAGGTGATTTGAAAGACAGGAGAGTCTTACATTTTAAAGTGAAGTTAATAGC 2613  
 2281 TTGAAAGAGAGATTAATAATGATATGACCATGCTAATAGAGGTGATCTGTTGTTA 2360  
 2614 TTGAAAGAGAGATTAATAATGATATGACCATGCTAATAGAGGTGATCTGTTGTTA 2673  
 2341 GCACTGGAATTCCTATTTTCAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2400  
 2674 GCACTGGAATTCCTATTTTCAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2733  
 2401 AATATCATAG 2460  
 2734 AATATCATAG 2793  
 2461 TCTGTGATTTCAAAATTAACCAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2520  
 2794 TCTGTGATTTCAAAATTAACCAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2853  
 2521 AGTGTGCTTTCTCCCAACAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2580  
 2854 AGTGTGCTTTCTCCCAACAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2913  
 2581 CTGTGCTTTCAACAG 2640  
 2914 CTGTGCTTTCAACAG 2973  
 2641 CTTCCTAATGACAG 2700  
 2974 CTTCCTAATGACAG 3033  
 2701 CAGTTAATCATAG 2760  
 3034 CAGTTAATCATAG 3093  
 2761 GCAG 2820  
 3094 GCAG 3153  
 2821 AATGAAATTAATGACAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2880  
 3154 AATGAAATTAATGACAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 3213  
 2881 ATGTTTCTAATCGAGCTCTTAAATTTGATGGA 2913  
 3214 ATGTTTCTAATCGAGCTCTTAAATTTGATGGA 3246

RESULT 2  
 US-08-252-995D-3  
 Sequence 3, Application US/08252995D  
 Patent No. 5650501  
 GENERAL INFORMATION:  
 APPLICANT: Dennis, James W  
 APPLICANT: Heffernan, Mike  
 APPLICANT: Fode, Carol  
 TITLE OF INVENTION: NOVEL SERINE/THREONINE KINASE  
 NUMBER OF SEQUENCES: 14  
 CORRESPONDENCE ADDRESS:  
 ADDRESSEE: BERSEKIN & PARR  
 STREET: 40 King Street West  
 CITY: Toronto  
 STATE: Ontario  
 COUNTRY: Canada  
 ZIP: M5H 3Y2  
 COMPUTER READABLE FORM:  
 MEDIUM TYPE: Floppy disk  
 OPERATING SYSTEM: IBM PC compatible  
 SOFTWARE: Patent: PC-DOS/MS-DOS  
 CURRENT APPLICATION DATA:  
 APPLICATION NUMBER: US/08/252,995D  
 FILING DATE: 02-JUN-1994  
 CLASSIFICATION: 536

ATTORNEY/AGENT INFORMATION:  
 NAME: Kurydyk, Linda M  
 REGISTRATION NUMBER: 34,971  
 REFERENCE/DOCKET NUMBER: 3153-96  
 TELECOMMUNICATION INFORMATION:  
 TELEPHONE: (416) 364-7311  
 TELEFAX: (416) 361-1398  
 INFORMATION FOR SEQ ID NO: 3:  
 SEQUENCE CHARACTERISTICS:  
 LENGTH: 3447 base pairs  
 TYPE: nucleic acid  
 STRANDEDNESS: single  
 TOPOLOGY: linear  
 MOLECULE TYPE: cDNA  
 ORIGINAL SOURCE:  
 ORGANISM: Mus musculus  
 DEVELOPMENTAL STAGE: Lymphoid cDNA library  
 IMMEDIATE SOURCE:  
 LIBRARY: Murine Lymphoid  
 CLONE: MGA-resistant chop clones  
 FEATURE:  
 NAME/KEY: 5'UTR  
 LOCATION: 1..205  
 FEATURE:  
 NAME/KEY: CDS  
 LOCATION: 206..2980  
 FEATURE:  
 NAME/KEY: 3'UTR  
 LOCATION: 2981..3447  
 US-08-252-995D-3

Query Match 64.5% Score 1879; DB 1; Length 3447;  
 Best Local Similarity 80.9%; Pred. No. 0;  
 Matches 2362; Conservative 0; Mismatches 410; Indels 147; Gaps 9;  
 1 ATGGGACCTGTCATCGGGGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 60  
 206 ATGGGAG 265  
 61 GATCAATTTGCTGTGTCATGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 120  
 266 GATCAATTTGCTGTGTCATGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 325  
 121 AATATGATGATGAG 180  
 326 AATATGATGATGAG 385  
 181 GTGAAATTAATGAG 240  
 386 GTGAAATTAATGAG 445  
 241 GATGCAATTAATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 300  
 446 GATGCAATTAATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 505  
 301 CTAAAGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 360  
 506 CTAAAGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATG 565  
 361 ATCAAG 420  
 566 ATCAAG 625  
 421 AACCTCTAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 480  
 626 AACCTCTAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 685  
 481 CTGAAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 540  
 686 CTGAAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 745  
 541 GAAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 600

746 GAATTCGAACTCGAAGTGCATGGACTTGATCTGATATTTGGTCACTTGGGCTGATG 805.  
QY TTTTATACATTAATTATCGGAGAGACCACTTTCGACATGACACAGTCAAGAACATTA 660  
Db TCTTATACCTTACTTATGGAAGACCACTTTTGAACATGACACAGTCAAGAACATTA 865  
QY AATAAGATGATGCGAATTAATGAAGCCACTTTTGGTCAATAGAGCGAAGAC 720  
Db AACAAAGATGCTCGGCAATTAATGAAGCCACTTTTGGTCAATAGAGCGAAGAC 925  
QY CTATATCACAGTACTTCTGATGAAGCCCTGAGATCGGTTAAGTGTCTTCTGATG 780  
Db CTATATCACAGTACTTCTGATGAAGCCCTGAGATCGGTTAAGTGTCTTCTGATG 985  
QY GACCATCTTTTATGTCCTCCGAAATCTTCAACAAAAGTAAAGATTAGAACTGTGAA 840  
Db GACCATCTTTTATGTCCTCCGAAATCTTCAACAAAAGTAAAGATTAGAACTGTGAA 1045  
QY GACTCAATTAATGAGGAGGAGTCCCAATTTCTACTGCAATTAACAGCTTCCAGTACC 900  
Db GACTCAATTAATGAGGAGGAGTCCCAATTTCTACTGCAATTAACAGCTTCCAGTACC 1105  
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Db AGTATAGATGATGATTTATTTGACAAAGAGAGCTTTGATGATGATGATGATGAT 1162  
QY AAAATGACTGATTTTCCAAAGAAATTAAGTCTCACTGATTTTCTTCAAGAGATGA 1020  
Db AAAATGACTGATTTTCCAAAGAAATTAAGTCTCACTGATTTTCTTCAAGAGATGA 1219  
QY AAACATTTTATCTAGTGGGGAAT-----CAAGAAACAGTAAATGATGAGGGA 1074  
Db AAACATTTTATCTAGTGGGGAAT-----CAAGAAACAGTAAATGATGAGGGA 1279  
QY AGAGTAAATTCAGATGAGAAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1134  
Db AGAGTAAATTCAGATGAGAAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1339  
QY TCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1194  
Db TCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1396  
QY TGTCTAGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 1254  
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QY AGTACTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1374  
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QY TGAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1614  
Db TGAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1708  
QY CAGCAAAATACATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1674  
Db CAGCAAAATACATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1768

QY GAATGTTTTTGGCTCAGATCTCTTTCTGAAACAGACAGACATGAGGGATGAGACCA 1734  
Db GA-----GCCGGGCTCATCTCTCATTTCTGAAACAAACAGAAATGATGAGGCTG 1822  
QY CAGTGGGATTAACAAATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1794  
Db CAGTGGGATTAACAAATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1882  
QY TTAACCAATCAGACAGAAACCAAAAGGCTGTGAGACATCTTGAATCAAGAGAG 1854  
Db TTAACCAATCAGACAGAAACCAAAAGGCTGTGAGACATCTTGAATCAAGAGAG 1942  
QY GTGTGTGAGGCTTGTAAAGGATGATGATGATGATGATGATGATGATGATGATGAT 1914  
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QY ATATCTAGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1974  
Db ATATCTAGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2062  
QY CTGCTGATGACACACCTCTCACTGACACACATGATGATGATGATGATGATGATGAT 2034  
Db CTGCTGATGACACACCTCTCTGCTGATGACACACATGATGATGATGATGATGATGAT 2122  
QY CCAAGAAATTAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 2094  
Db CCAAGAAATTAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 2182  
QY AAATCTCCCAAAATACATTAATTAACAGATGATGATGATGATGATGATGATGATGAT 2154  
Db AAATCTCCCAAAATACATTAATTAACAGATGATGATGATGATGATGATGATGATGAT 2242  
QY CTTGCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2214  
Db CTTGCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2302  
QY TTTTATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTA 2274  
Db TTTTATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTA 2362  
QY AATGCTGAAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 2334  
Db AATGCTGAAAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAGGAG 2422  
QY TGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2394  
Db TGTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2482  
QY TTTCCCAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 2454  
Db TTTCCCAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 2542  
QY CTTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2514  
Db CTTCTGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2599  
QY ATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2574  
Db ATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2660  
QY GATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2634  
Db GATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2708  
QY GATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2694  
Db GATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 2764  
QY GCTACAGATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 2754  
Db GCTACAGATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAATTAAT 2824

QY 2755 GTCAGGAGAGAGTCTTCTATCACTTAACTACCAATAGTCAACAACAGTAT 2814  
DB 2825 GTCAGGAGAGAGTCTTCTATCACTTAACTACCAATAGTCAACAACAGTAT 2884  
QY 2815 GAGAAATGAAATTAACCAAGTCACTAATCAAGAAATTAAGTCTTCTATC 2874  
DB 2885 GAGAAATGAAATTAACCAAGTCACTAATCAAGAAATTAAGTCTTCTATC 2944  
QY 2875 CTTTGAATGTTTCTAATCCGACTCTAATTTTCTTGA 2913  
DB 2945 CTTTGAATGTTTCTAATCCGACTCTAATTTTCTTGA 2983

US-08-834-108-3  
Sequence 3, Application US/08834108

GENERAL INFORMATION:  
PATENT NO. 5976893  
APPLICANT: Dennis, James W  
APPLICANT: Heffernan, Mike  
APPLICANT: Rode, Carol  
TITLE OF INVENTION: NOVEL SERINE/THREONINE KINASE  
NUMBER OF SEQUENCES: 14  
CORRESPONDENCE ADDRESS:  
ADDRESS: BERSKIN & PARR  
STREET: 40 King Street West  
CITY: Toronto  
STATE: Ontario  
COUNTRY: Canada  
ZIP: M5H 3Y2  
COMPUTER READABLE FORM:  
MEDIUM TYPE: Floppy disk  
OPERATING SYSTEM: PC-DOS/MS-DOS  
SOFTWARE: Patent Release #1.0, Version #1.30  
CURRENT APPLICATION DATA:  
APPLICATION NUMBER: US/08/834,108  
FILING DATE:  
CLASSIFICATION: 536  
ATTORNEY/AGENT INFORMATION:  
NAME: Kurydyk, Linda M  
REGISTRATION NUMBER: 34,971  
REFERENCE/DOCKET NUMBER: 3153-210  
TELECOMMUNICATION INFORMATION:  
TELEPHONE: (416) 364-7311  
TELEFAX: (416) 361-1398  
INFORMATION FOR SEQ ID NO: 3:  
SEQUENCE CHARACTERISTICS:  
LENGTH: 3447 base pairs  
TYPE: nucleic acid  
STRANDEDNESS: single  
TOPOLOGY: linear  
MOLECULAR TYPE: cDNA  
ORIGINAL SOURCE:  
ORGANISM: Mus musculus  
DEVELOPMENTAL STAGE: Lymphoid cDNA Library  
IMMEDIATE SOURCE: Lymphoid  
LIBRARY: Murine Lymphoid  
CLONE: WGA-resistant chop clones  
FEATURE:  
NAME/KEY: 5' UTR  
LOCATION: 1..205  
FEATURE:  
NAME/KEY: CDS  
LOCATION: 206..2980  
FEATURE:  
NAME/KEY: 3' UTR  
LOCATION: 2981..3447  
US-08-834-108-3

Query Match 64.5%; Score 1879; DB 2; Length 3447;  
Best Local Similarity 80.9%; Pred. No. 0;  
Matches 2362; Conservative 0; Mismatches 410; Indels 147; Gaps 9;

QY 1 ATGGCCACTGATCGGGGAGAAATCGAGATTTTAAAGTTGAAATCTGCTTGTTAA 60  
DB 206 ATGGCCGCGTGATCGGGGAGAGATCGAGATTTTAAAGTTGAAATCTGCTTGTTAA 265  
QY 61 GATTCATTTTCTGATGTTTCAAGAGTGAATCTTCACTGCTTGAAGTGCATC 120  
DB 266 GATTCATTTTCTGATGTTTCAAGAGTGAATCTTCACTGCTTGAAGTGCATC 325  
QY 121 AAAATGATTAAGTAAAGGATGTAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 180  
DB 326 AAAATGATTAAGTAAAGGATGTAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 385  
QY 181 GTGAAATACATTCGCAATGAAACATCTTCTAATCTTGAAGTTTAACTATTGTA 240  
DB 386 GTGAAATACATTCGCAATGAAACATCTTCTAATCTTGAAGTTTAACTATTGTA 445  
QY 241 GATGCAATTAATGTTTCTGTTTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGA 300  
DB 446 GATGCAATTAATGTTTCTGTTTGAAGTGAAGTGAAGTGAAGTGAAGTGAAGTGA 505  
QY 301 CTAAAGATTAAGTAAACCTTCTGAGAAATGAGTGCACCTTCATGACCAAGATC 360  
DB 506 CTAAAGATTAAGTAAACCTTCTGAGAAATGAGTGCACCTTCATGACCAAGATC 565  
QY 361 ATCAAGAGATGTTGATCTTCAATCTGATGATATCAACCGGACTCACTTTCT 420  
DB 566 ATCAAGAGATGTTGATCTTCAATCTGATGATATCAACCGGACTCACTTTCT 625  
QY 421 AACTCTGATGATCTGATATGATGATGATGATGATGATGATGATGATGATGAT 480  
DB 626 AACTCTGATGATCTGATATGATGATGATGATGATGATGATGATGATGATGAT 685  
QY 481 CTGAAATGAGGACATGAAAGGACATATGATGATGATGATGATGATGATGATGAT 540  
DB 686 CTGAAATGAGGACATGAAAGGACATATGATGATGATGATGATGATGATGATGAT 745  
QY 541 GAAATGAGGACATGAAAGGACATATGATGATGATGATGATGATGATGATGATGAT 600  
DB 746 GAAATGAGGACATGAAAGGACATATGATGATGATGATGATGATGATGATGATGAT 805  
QY 601 TTTTATACATTAATCTTATGAGGAGACACCTTCTGACATGACATGACATGACAT 660  
DB 806 TTTTATACATTAATCTTATGAGGAGACACCTTCTGACATGACATGACATGACAT 865  
QY 661 AATAAGTATGATGAGGACATATGATGATGATGATGATGATGATGATGATGAT 720  
DB 866 AATAAGTATGATGAGGACATATGATGATGATGATGATGATGATGATGATGAT 925  
QY 721 CTTATTCACAGTATCTTGTGAGAAATCCAGAGATCGTTAAGTCTTCTGATGAT 780  
DB 926 CTTATTCACAGTATCTTGTGAGAAATCCAGAGATCGTTAAGTCTTCTGATGAT 985  
QY 781 GACCATCTTTTATGATGATGATGATGATGATGATGATGATGATGATGATGAT 840  
DB 986 GACCATCTTTTATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1045  
QY 841 GACTCAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 900  
DB 1046 GACTCAATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1105  
QY 901 AGTATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 960  
DB 1106 AGTATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1162  
QY 961 AAAATGATTAAGTAAAGGATGTAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1020  
DB 1163 AAAATGATTAAGTAAAGGATGTAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1219  
QY 1021 AACTGTTTATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1074  
DB 1220 AACTGTTTATGATGATGATGATGATGATGATGATGATGATGATGATGATGATGAT 1279